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SEQUENCE LISTING

(1) GENERAL INFORMATION

- (i) APPLICANT: BILLING-MEDEL, PATRICIA
COHEN, MAURICE
COLPITTS, TRACEY L.
FRIEDMAN, PAULA N.
GORDON, JULIAN
GRANADOS, EDWARD N.
HODGES, STEVEN C.
KLASS, MICHAEL R.
KRATOCHVIL, JON D.
ROBERTS-RAPP, LISA
RUSSELL, JOHN C.
STROUPE, STEPHEN D.
YU, HONG
- (ii) TITLE OF THE INVENTION: REAGENTS AND METHODS USEFUL
FOR DETECTING DISEASES OF THE URINARY TRACT
- (iii) NUMBER OF SEQUENCES: 22
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: Abbott Laboratories
 - (B) STREET: 100 Abbott Park Road
 - (C) CITY: Abbott Park
 - (D) STATE: IL
 - (E) COUNTRY: USA
 - (F) ZIP: 60064-3500
- (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Diskette
 - (B) COMPUTER: IBM Compatible
 - (C) OPERATING SYSTEM: DOS
 - (D) SOFTWARE: FastSEQ for Windows Version 2.0
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER:
 - (B) FILING DATE:
 - (C) CLASSIFICATION:
- (vii) PRIOR APPLICATION DATA:
 - (A) APPLICATION NUMBER: 08/869,579
 - (B) FILING DATE: 05-JUN-1997
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Becker, Cheryl L.
 - (B) REGISTRATION NUMBER: 35,441
 - (C) REFERENCE/DOCKET NUMBER: 6107.US.P1
- (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: 847/935-1729
 - (B) TELEFAX: 847/938-2623
 - (C) TELEX:

(2) INFORMATION FOR SEQ ID NO:1:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 196 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

CTCCACTGCA	ACCACCCAGA	GCCATGGCTC	CCCAGAGCTG	CATCGTAGCT	GTCTTTGCCA	60
TTTTCTGCAT	CTCCAGGCTC	CTCTGCTCAC	ACGGAGCCCC	AGTGGCCCCC	ATGACTCCTT	120
ACCTGATGCT	GTGCCAGCCA	CACAAGAGAT	GTGGGGACAA	GTTCTACGAC	CCCCTGCAGC	180
ACTGTTGCTA	TGATGA					196

(2) INFORMATION FOR SEQ ID NO:2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 236 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

TTCTACGACC	CCCTGCAGCA	CTGTTGCTAT	GATGATGCCG	TCGTGCCCTT	GGCCAGGACC	60
CAGACGTGTG	GAAACTGCAC	CTTCAGAGTC	TGCTTTGAGC	AGTGCTGCCC	CTGGACCTTC	120
ATGGTGAAGC	TGATAAACCA	GAAGTGCAGC	TCAGCCCCGA	CCTCGGATGA	CAGGCTTTGT	180
CGCAGTGTCA	GCTAATGGAA	CATCAGGGGA	ACGATGACTC	CTGGATTCTC	CTTCCT	236

(2) INFORMATION FOR SEQ ID NO:3:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 417 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

ATAAAGTACA	TAAATTTATT	ATGATCCTGT	GCTTTGCTCC	TGGATAGAAG	AGAACTTGAG	60
ATGATGGAAA	ATCTCACAGA	AGGGTGAAGG	GTTCCCCAGC	CAGCCTTGCG	GTGGTGCCGG	120
GTTGATGTTG	GCCATAGGGG	TCATCAGGAG	GGTTCAGAGT	CCCTGGGCCC	CCATCAGCAC	180
ACGCATCTCA	TGCTGTCCAG	AATTCTAGAA	GGGGACCTGT	GGGCCTCCTC	AGGGCAGCGG	240
CAGGGTGGGC	CTCAGACAGG	TCACAGAATG	AAGTGGGCAG	TTGAGTGTGT	GTTTCTCTGG	300
CCCCCAAACA	GCCACTCAGC	ATCCCAGATC	TCAGGTAACA	CCAGCCTCTT	TCTCCAGGCC	360
CACCCAGGAA	GGAGAATCCA	GGAGTCATCG	TTCCCCTGAT	GTTCCATTAG	CTGACAC	417

(2) INFORMATION FOR SEQ ID NO:4:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 763 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CTCCACTGCA	ACCACCCAGA	GCCATGGCTC	CCCAGAGCTG	CATCGTAGCT	GTCTTTGCCA	60
TTTTCTGCAT	CTCCAGGCTC	CTCTGCTCAC	ACGGAGCCCC	AGTGGCCCCC	ATGACTCCTT	120
ACCTGATGCT	GTGCCAGCCA	CACAAGAGAT	GTGGGGACAA	GTTCTACGAC	CCCCTGCAGC	180
ACTGTTGCTA	TGATGATGCC	GTCGTGCCCT	TGGCCAGGAC	CCAGACGTGT	GGAAACTGCA	240

CCTTCAGAGT	CTGCTTTGAG	CAGTGCTGCC	CCTGGACCTT	CATGGTGAAG	CTGATAAACC	300
AGAACTGCGA	CTCAGCCCGG	ACCTCGGATG	ACAGGCTTTG	TGCGAGTGTC	AGCTAATGGA	360
ACATCAGGGG	AACGATGACT	CCTGGATTCT	CCTTCCTGGG	TGGGCCTGGA	GAAAGAGGCT	420
GGTGTTACCT	GAGATCTGGG	ATGCTGAGTG	GCTGTTTGGG	GGCCAGAGAA	ACACACACTC	480
AACTGCCCCAC	TTCATTCTGT	GACCTGTCTG	AGGCCACCCC	TGCCGCTGCC	CTGAGGAGGC	540
CCACAGGTCC	CCTTCTAGAA	TTCTGGACAG	CATGAGATGC	GTGTGCTGAT	GGGGGCCCAG	600
GGACTCTGAA	CCCTCCTGAT	GACCCGTATG	GCCAACATCA	ACCCGGCACC	ACCCCAAGGC	660
TGGCTGGGGA	ACCCTTCACC	CTTCTGTGAG	ATTTTCCATC	ATCTCAAGTT	CTCTTCTATC	720
CAGGAGCAAA	GCACAGGATC	ATAATAAATT	TATGTACTTT	ATA		763

(2) INFORMATION FOR SEQ ID NO:5:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 763 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

CTCCACTGCA	ACCACCCAGA	GCCATGGCTC	CCCGAGGCTG	CATCGTAGCT	GTCTTTGCCA	60
TTTTCTGCAT	CTCAGGCTC	CTCTGCTCAC	ACGGAGCCCC	AGTGGCCCCC	ATGACTCCTT	120
ACCTGATGCT	GTGCCAGCCA	CACAAGAGAT	GTGGGGACAA	GTTCTACGAC	CCCCTGCAGC	180
ACTGTTGCTA	TGATGATGCC	GTCGTGCCCT	TGGCCAGGAC	CCAGACGTGT	GGAAACTGCA	240
CCTTCAGAGT	CTGCTTTGAG	CAGTGCTGCC	CCTGGACCTT	CATGGTGAAG	CTGATAAACC	300
AGAACTGCGA	CTCAGCCCGG	ACCTCGGATG	ACAGGCTTTG	TGCGAGTGTC	AGCTAATGGA	360
ACATCAGGGG	AACGATGACT	CCTGGATTCT	CCTTCCTGGG	TGGGCCTGGA	GAAAGAGGCT	420
GGTGTTACCT	GAGATCTGGG	ATGCTGAGTG	GCTGTTTGGG	GGCCAGAGAA	ACACACACTC	480
AACTGCCCCAC	TTCATTCTGT	GACCTGTCTG	AGGCCACCCC	TGCCGCTGCC	CTGAGGAGGC	540
CCACAGGTCC	CCTTCTAGAA	TTCTGGACAG	CATGAGATGC	GTGTGCTGAT	GGGGGCCCAG	600
GGACTCTGAA	CCCTCCTGAT	GACCCSTATG	GCCAACATCA	ACCCGGCACC	ACCCCAAGGC	660
TGGCTGGGGA	ACCCTTCACC	CTTCTGTGAG	ATTTTCCATC	ATCTCAAGTT	CTCTTCTATC	720
CAGGAGCAAA	GCACAGGATC	ATAATAAATT	TATGTACTTT	ATA		763

(2) INFORMATION FOR SEQ ID NO:6:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 68 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

AGCTCGGAAT	TCCGAGCTTG	GATCCTCTAG	AGCGGCCGCC	GACTAGTGAG	CTCGTCGACC	60
CGGGAATT						68

(2) INFORMATION FOR SEQ ID NO:7:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 68 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

AATTAATTCC	CGGGTCGACG	AGCTCACTAG	TCGGCGGCCG	CTCTAGAGGA	TCCAAGCTCG	60
GAATTCCC						68

(2) INFORMATION FOR SEQ ID NO:8:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 24 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

AGCGGATAAC AATTTACAC AGGA

24

(2) INFORMATION FOR SEQ ID NO:9:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

TGTAAACGA CGGCCAGT

18

(2) INFORMATION FOR SEQ ID NO:10:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

TGGAAACTGC ACCTTCAG

18

(2) INFORMATION FOR SEQ ID NO:11:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 20 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

CAAAGCAGAC TCTGAAGGTG

20

(2) INFORMATION FOR SEQ ID NO:12:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 18 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

ACGCATCTCA TGCTGTCC

18

(2) INFORMATION FOR SEQ ID NO:13:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs

Arg Cys Gly Asp Lys Phe Tyr Asp Pro Leu Gln His Cys Cys Tyr Asp
 50 55 60
 Asp Ala Val Val Pro Leu Ala Arg Thr Gln Thr Cys Gly Asn Cys Thr
 65 70 75 80
 Phe Arg Val Cys Phe Glu Gln Cys Cys Pro Trp Thr Phe Met Val Lys
 85 90 95
 Leu Ile Asn Gln Asn Cys Asp Ser Ala Arg Thr Ser Asp Asp Arg Leu
 100 105 110
 Cys Arg Ser Val Ser
 115

(2) INFORMATION FOR SEQ ID NO:18:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 32 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: None

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

Pro Leu Gln Pro Pro Arg Ala Met Ala Pro Arg Gly Cys Ile Val Ala
 1 5 10 15
 Val Phe Ala Ile Phe Cys Ile Ser Arg Leu Leu Cys Ser His Gly Ala
 20 25 30

(2) INFORMATION FOR SEQ ID NO:19:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 30 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: None

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Leu Cys Gln Pro His Lys Arg Cys Gly Asp Lys Phe Tyr Asp Pro Leu
 1 5 10 15
 Gln His Cys Cys Tyr Asp Asp Ala Val Val Pro Leu Ala Arg
 20 25 30

(2) INFORMATION FOR SEQ ID NO:20:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 31 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: None

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

Thr Gln Thr Cys Gly Asn Cys Thr Phe Arg Val Cys Phe Glu Gln Cys
 1 5 10 15
 Cys Pro Trp Thr Phe Met Val Lys Leu Ile Asn Gln Asn Cys Asp
 20 25 30

(2) INFORMATION FOR SEQ ID NO:21:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 8 amino acids

- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

AGGCTGCATC GTAGCTGTCT TT

22

(2) INFORMATION FOR SEQ ID NO:14:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 22 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

CGCATCTCAT GCTGTCCAGA AT

22

(2) INFORMATION FOR SEQ ID NO:15:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 23 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

CCCAGTCACG ACGTTGTAAA ACG

23

(2) INFORMATION FOR SEQ ID NO:16:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 26 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

GCGGCCGCGG CTGACACTGC GACAAA

26

(2) INFORMATION FOR SEQ ID NO:17:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 117 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: None

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

Pro	Leu	Gln	Pro	Pro	Arg	Ala	Met	Ala	Pro	Arg	Gly	Cys	Ile	Val	Ala	
1				5				10						15		
Val	Phe	Ala	Ile	Phe	Cys	Ile	Ser	Arg	Leu	Leu	Cys	Ser	His	Gly	Ala	
		20					25						30			
Pro	Val	Ala	Pro	Met	Thr	Pro	Tyr	Leu	Met	Leu	Cys	Gln	Pro	His	Lys	
		35				40						45				

- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

Asp Tyr Lys Asp Asp Asp Asp Lys
1 5

(2) INFORMATION FOR SEQ ID NO:22:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 21 amino acids
- (B) TYPE: amino acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu Asn Met His Thr Glu His
1 5 10 15
His His His His His
20